

Figure 1

ATF1:	QTTKTDDPQ-LKREIRLMKNREAARECR-RKKK	(205 - 235 of SEQ ID NO:1)
CREB:	PTQPAEE-AARKREVRLMKNREAARECR-RKKK	(275 - 305 of SEQ ID NO:2)
GCN4:	IDDPAALKR-ARNTEAARRSRARKL-	(224 - 247 of SEQ ID NO:3)

Figure 2

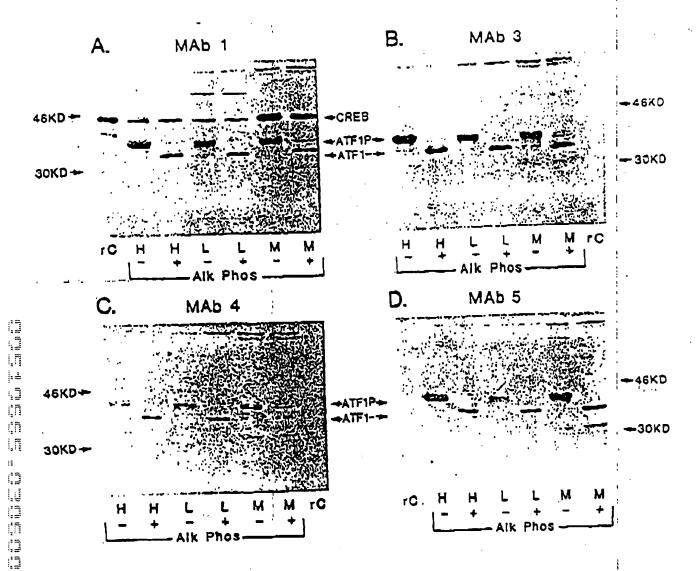
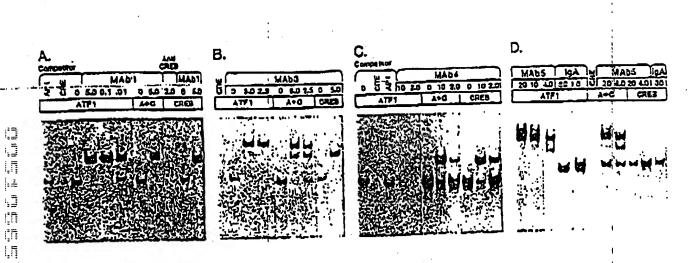
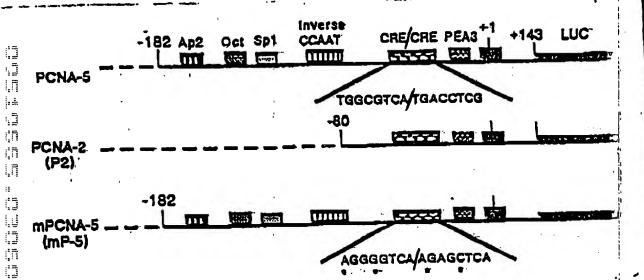
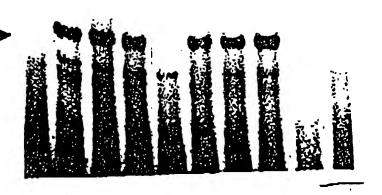


FIGURE 3



PCNA-LUC constructs





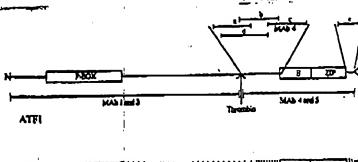


FIGURE 7

MAD1 MAD3 MAD4 MAD5

-46KD
-30KD
-30KD
-21.5KD

C-terminal
fragment
Thrombin
digest (h)

MAD3 MAD4 MAD5

-46KD
-30KD
-21.5KD

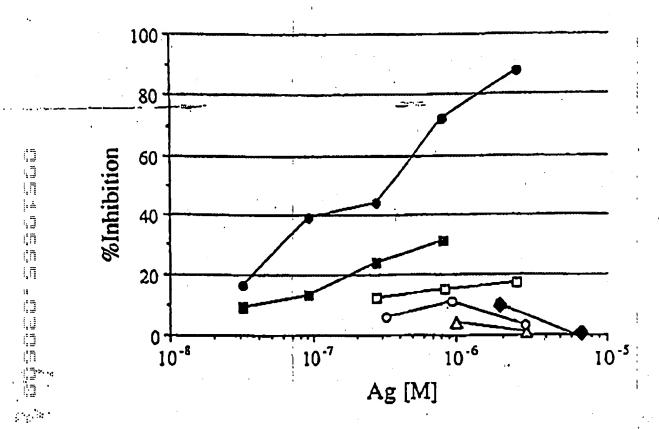
(n (n (n

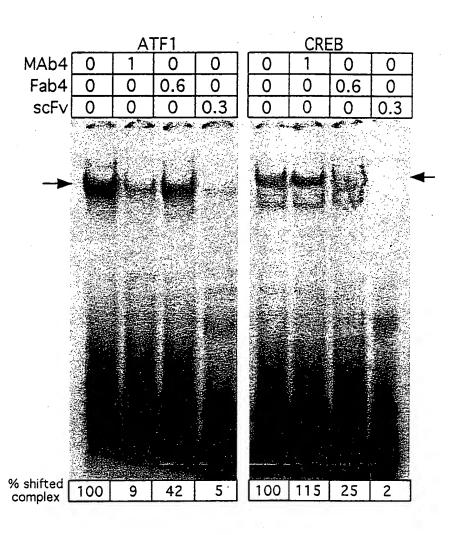
FIGURE 8

 $A(\widehat{\mathbb{Q}})$

M1 M3 M4 M5

FIGURE 9





Figurell

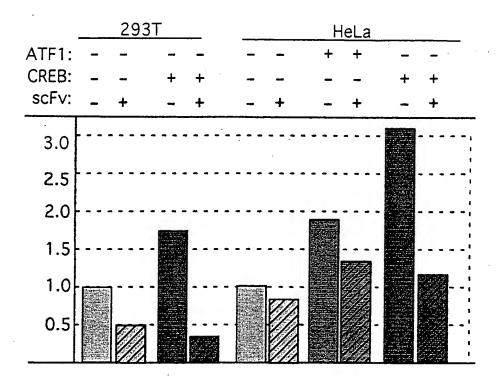


Figure 12

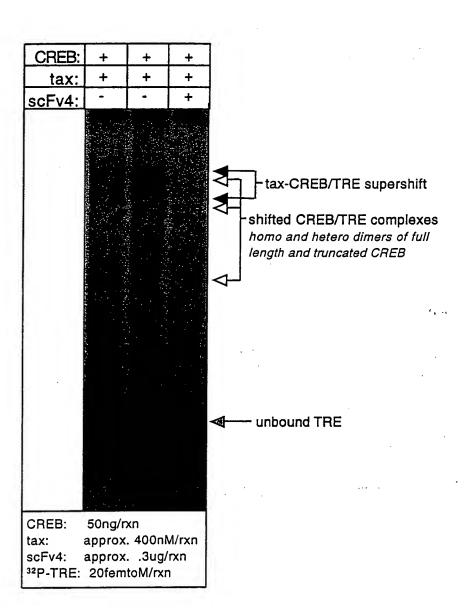


Figure 13